Applicant: Walhout et al. Attorney's Docket No.: 07917-0232US1 / UMMS 03-137; DFCI 940

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Amendments to the Claims

This listing of claims replaces all prior versions and listings of claims in the application.

<u>Listing of Claims</u>:

1. (Currently amended) A method of identifying a protein that binds to a bait element, the method comprising:

obtaining a <u>yeast or mammalian</u> cell whose genome comprises one or more integrated bait-reporter constructs, wherein each of the one or more bait-reporter constructs comprises

- (a) a single copy of a bait element flanked by lambda recombination sites, wherein the bait element comprises at least 250 base pairs; and
 - (b) a reporter gene;

transforming the cell with an expression vector encoding a fusion protein comprising an activation domain; and

assessing activation of the reporter gene,

wherein activation of the reporter gene indicates that the activation domain fusion protein has bound to the bait element.

- 2. (Original) The method of claim 1, wherein the cell is a yeast cell.
- 3. (Original) The method of claim 2, wherein the yeast cell is a YM4271 cell.
- 4. (Original) The method of claim 2, wherein the reporter gene is HIS3.
- 5. (Original) The method of claim 2, wherein the reporter gene is LacZ.
- 6. (Original) The method of claim 1, wherein two reporter genes are used.

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7. (Original) The method of claim 6, wherein the cell is a yeast cell and the reporter genes are HIS3 and LacZ.

- 8. (Original) The method of claim 1, wherein the cell is a mammalian cell.
- 9. (Original) The method of claim 8, wherein the reporter gene is a fluorescent gene.
- 10. (Original) The method of claim 9, wherein the fluorescent gene is selected from the group consisting of luciferase, green fluorescent protein, yellow fluorescent protein, red fluorescent protein, blue fluorescent protein, or a fluorescent variant of any of said proteins.
- 11. (Original) The method of claim 1, wherein the bait element comprises more than 500 base pairs.
- 12. (Original) The method of claim 1, wherein the fusion protein is an activation domain-transcription factor fusion protein.
- 13. (Currently amended) The method of claim [[1]] 2, wherein assessing activation comprises plating the yeast cell on nutrient deficient plates under growth conditions, and observing any growth of yeast cells.
- 14. (Currently amended) The method of claim [[1]] 2, wherein assessing activation comprises determining the color of the yeast cells.
- 15. (Original) The method of claim 1, further comprising selecting a cell in which a reporter gene is activated; and isolating the cDNA encoding an activation domain fusion protein that is bound to the bait element from the cell.

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16. (Original) The method of claim 15, further comprising determining the sequence of the isolated cDNA.

17. (Original) The method of claim 1, further comprising contacting the cell with a test compound, and evaluating the effect of the test compound on binding of the bait element to the activation domain fusion protein.